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OCT 0 3 2001

TECH CENTER 1600/2900

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SUGANUMA, MASASHI
KAWABE, TAKUMI
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15 1 5 10 ⊴210⇒ 58 4211 × 11 -1112 - PRT -113: Artificial Sequence

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- 224

+22: Description of Artificial Sequence: Synthetic peptide

· 400 · 87

Leu Tyr Thr Ser Pro Ser Asn Phe Glu Asn Leu
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· 210 · 88

-211 - 11 -212 - PRT

·213: Artificial Sequence

< 220 h

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4312 - PRT

«213 · Artificial Sequence

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1 5 10

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peptide <400 - 381 Leu Phe Ser Ser Pro Ser Asn Trp Glu Asn Leu 1 Ε, 3210 - 282 0211 - 11 3212 - PRT <213 - Artificial Sequence 4120 × <223 - Description of Artificial Sequence: Synthetic peptide <400 - 182 Lou The Ser Ser Pro Ser Gln Pro Glu Asn Leu 5 <210 / 283 <211 × 11 <212 · FRT <213 - Artificial Sequence <223 - Description of Artificial Sequence: Synthetic</p> reptide <400 > 283 Leu Phe Ser Ser Pro Ser Gln Phe Glu Asn Leu 1 5 <210> 284 <211> 11 <212> FRT <213: Artificial Sequence <223> Description of Artificial Sequence: Synthetic peptide . 400 - 294

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-2.1> 11
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```
peptide
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+ 2.12 + PRT
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peptide
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Leu Tyr Arg Ser Pro Ser Gly Pro Glu Asn Leu
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<400> 1863
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Leu Tyr Arg Ser Pro Ser Ile Pro Glu Asn Leu
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<210> 1865
<211> 11
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<400> 1865
Leu Tyr Arg Ser Pro Ser Leu Pro Glu Asn Leu
<210> 1866
<211> 11
<212> PRT
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<223> Description of Artificial Sequence: Synthetic
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Leu Tyr Arg Ser Pro Ser Lys Pro Glu Asn Leu
<2105 1867
4.2110-11
<2212> PRT
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<400> 1867
Leu Tyr Arg Ser Pro Ser Met Pro Glu Asn Leu
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<400> 1868
Leu Tyr Arg Ser Pro Ser Phe Pro Glu Asn Leu
<210> 1869
<211> 11
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<400> 1869
Leu Tyr Arg Ser Pro Ser Pro Pro Glu Asn Leu
<210> 1870
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<212> PRT
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Leu Tyr Arg Ser Pro Ser Ser Pro Glu Asn Leu
<210> 1871
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<4(0)> 1871
Leu Tyr Arg Ser Pro Ser Thr Pro Glu Asn Leu
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Leu Tyr Arg Ser Pro Ser Val Pro Glu Asn Leu
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peptide

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Leu Tyr Arg Ser Pro Ser Met Asp Glu Asn Leu
<210> 1879
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Leu Tyr Arg Ser Pro Ser Met Cys Glu Asn Leu
<210> 1880
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<400> 1880
Leu Tyr Arg Ser Pro Ser Met Gln Glu Asn Leu
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<210> 1881
<211> 11
<212> PRT
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<400> 1881
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<210> 1882
<211> 11
<212> PRT
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<400> 1882
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                5
<210> 1883
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<212> PRT
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<:i00> 1883
Leu Tyr Arg Ser Pro Ser Met His Glu Asn Leu
 1 5
<210> 1884
<211> 11
<212> PRT
<213> Artificial Sequence
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<210> 1885
<211> 11
<212> PRT
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Leu Tyr Arg Ser Pro Ser Met Leu Glu Asn Leu
<210> 1886
<211> 11
<212> PRT
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<400> 1886
Leu Tyr Arg Ser Pro Ser Met Lys Glu Asn Leu
<210> 1887
<211> 11
<212> PRT
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<400> 1887
Leu Tyr Arg Ser Pro Ser Met Met Glu Asn Leu
<210> 1888
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<212> PRT
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<223> Description of Artificial Sequence: Synthetic
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peptide

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<210> 1889
<211> 11
<212> PRT
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<400> 1889
Leu Tyr Arg Ser Pro Ser Met Pro Glu Asn Leu
<210> 1890
<211> 11
<212> PRT
<213> Artificial Sequence
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<400> 1890
Leu Tyr Arg Ser Pro Ser Met Ser Glu Asn Leu
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<110> 1891
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<400> 1892
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<212> PRT
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<210> 1894
<211> 11
<212> PRT
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      peptide
<400> 1894
Leu Tyr Arg Ser Pro Ser Met Val Glu Asn Leu
<210> 1895
<211> 9
<212> PRT
<213> Artificial Sequence
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<400> 1895
Tyr Gly Gly Pro Gly Gly Gly Asn
<210> 1896
<111> 11
<1112 > PRT
<2233 Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
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<:400> 1896
Leu Ala Arg Ser Ala Ser Met Pro Glu Ala Leu
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<213> Artificial Sequence
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< 100> 1897
Leu Tyr Arg Ser Pro Ala Met Pro Glu Asn Leu
<210> 1898
<211> 8
<212> PRT
<213> Artificial Sequence
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<400> 1898
Trp Tyr Arg Ser Pro Ser Tyr Tyr
<210> 1899
<211> 11
<212> PRT
<213> Artificial Sequence
<220>
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<400> 1899
Tyr Gly Arg Lys Lys Arg Arg Gln Arg Arg Arg
<210> 1900
<211> 2055
<212> DNA
<113> Homo sapiens
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taastacete ettteeeeta gattteettt eattetgete aagtettege etgtgteega 120
tecetateta etttetetee tettgtagea ageeteagae teeaggettg agetaggttt 180
tgtttttctc ctggtgagaa ttcgaagacc atgtctacgg aactcttctc atccacaaga 240
gaggaaggaa getetggete aggaeceagt tttaggteta ateaaaggaa aatgttaaac 300
ctgctcctgg agagagacac ttcctttacc gtctgtccag atgtccctag aactccagtg 360
ggcaaatttc ttggtgattc tgcaaaccta agcattttgt ctggaggaac cccaaaatgt 420
tgcctcgatc tttcgaatct tagcagtggg gagataactg ccactcaget taccacttct 480
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tcagatgaat taatggagtt ttccctgaaa gatcaagaag caaaggtgag cagaagtggc 840
ctatateget eccegtegat gecagagaac ttgaacagge caagactgaa geaggtggaa 900
aaattcaagg acaacacaat accagataaa gttaaaaaaa agtatttttc tggccaagga 960
aageteagga agggettatg titaaagaag acagtetete tgtgtgacat tactateact 1020
cagatgetgg aggaagatte taaccagggg cacetgattg gtgattttte caaggtatgt 1080
gegetgecaa eegtgteagg gaaacaccaa gatetgaagt atgteaacce agaaacagtg 1140
gctgccttac tgtcggggaa gttccagggt ctgattgaga agttttatgt cattgattgt 1200
cgctatccat atgagtatct gggaggacac atccagggag ccttaaactt atatagtcag 1260
gaagaactgt ttaacttett tetgaagaag eccategtee etttggacae ecagaagaga 1320
ataatcatcg tgttccactg tgaattctcc tcagagaggg gcccccgaat gtgccgctgt 1380
ctgcgtgaag aggacaggte tetgaaceag tateetgeat tgtaetaeee agagetatat 1440
atcettaaag geggetaeag agaettettt reagaatata tggaactgtg tgaaccacag 1500
agetactgee etatgeatea teaggaceae aagaetgagt tgetgaggtg tegaageeag 1560
agcaaagtgc aggaagggga gcggcagctg cgggagcaga ttgcccttct ggtgaaggac 1620
atgageceat gataacatte cagecactgg etgetaacaa gteaceaaaa agacactgea 1680
gaaaccetga geagaaagag geettetgga tggeeaaace caagattatt aaaagatgte 1740
tetgeaaace aacaggetae caacttgtat ecaggeetgg gaatggatta ggttteagea 1800
gagetgaaag etggtggeag agteetggag etggetetat aaggeageet tgagttgeat 1860
agagatttgt attggttcag ggaactctgg cattcctttt cccaactcct catgtcttct 1920
cacaagccag ccaactettt etetetggge ttegggetat gcaagagegt tgtetacett 1980
ctttctttgt attttccttc tttgtttccc cctctttctt ttttaaaaat ggaaaaataa 2040
acactacaga atgag
<210> 1901
<211> 472
<212> PRT
<213> Homo sapiens
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Ser Gly Pro Ser Phe Arg Ser Asn Gln Arg Lys Met Leu Asn Leu Leu
Leu Glu Arg Asp Thr Ser Phe Thr Val Cys Pro Asp Val Pro Arg Thr
Pro Val Gly Lys Phe Leu Gly Asp Ser Ala Asn Leu Ser Ile Leu Ser
```

His Asp Gln His Leu Met Lys Cys Ser Pro Ala Gln Leu Leu Cys Ser 115 120 125

Gly Gly Thr Pro Lys Cys Cys Leu Asp Leu Ser Asn Leu Ser Ser Gly

Glu Ile Thr Ala Thr Gln Leu Thr Tnr Ser Ala Asp Leu Asp Glu Thr

Gly His Leu Asp Ser Ser Leu Gln Glu Val His Leu Ala Gly Met Asn
100 105 110

| 145 | | | | | 150 | | | | | 155 | | | | | 160 |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Glu | Met | Lys | Tyr | Leu
165 | Gly | Ser | Pro | Ile | Thr
170 | Thr | Val | Pro | Lys | Leu
175 | Asp |
| Lys | Asn | Pro | Asn
180 | Leu | Gly | Glu | Asp | Gln
185 | Ala | Glu | Glu | Ile | Ser
190 | Asp | Glu |
| Leu | Met | Glu
195 | Phe | Ser | Leu | Lys | Asp
200 | Gln | Glu | Ala | Lys | Val
205 | Ser | Arg | Ser |
| Gly | Leu
210 | Tyr | Arg | Ser | Pro | Ser
215 | Met | Pro | Glu | Asn | Leu
220 | Asn | Arg | Pro | Arç |
| Leu
225 | Lys | Gln | Val | Glu | Lys
230 | Phe | Lys | Asp | Asn | Thr
235 | Ile | Pro | Asp | Lys | Val
240 |
| Lys | Lys | Lys | Tyr | Phe
245 | Ser | Gly | Gln | Gly | Lys
250 | Leu | Arg | Lys | Gly | Leu
255 | Суя |
| Leu | Ľуs | Lys | Thr
260 | Val | Ser | Leu | Cys | Asp
265 | Il∈ | Thr | Ile | Thr | Gln
270 | Met | Let |
| Glu | Glu | Asp
275 | Ser | Asn | Gln | Gly | His
280 | Leu | Ile | Gly | Asp | Phe
285 | Ser | Lys | Val |
| Cys | Ala
290 | Leu | Pro | Thr | Val | Ser
295 | Gly | Lys | His | Gln | Asp
300 | Leu | Lys | Tyr | Val |
| Asn
305 | Pro | Glu | Thr | Val | Ala
310 | Ala | Leu | Leu | Ser | Gly
315 | Lys | Phe | Gln | Gly | Let
320 |
| Ile | Glu | Lys | Phe | Tyr
325 | Val | Ile | Asp | Cys | Arg
330 | Tyr | Pro | Tyr | Glu | Tyr
335 | Let |
| Gly | Gly | His | Ile
340 | Gln | Gly | Ala | Leu | Asn
345 | Leu | Tyr | Ser | Gln | Glu
350 | Glu | Leu |
| Phe | Asn | Phe
355 | Phe | Leu | Lys | Lys | Pro
360 | Ile | Val | Pro | Leu | Asp
365 | Thr | Gln | Lys |
| | Ile
370 | | Ile | | Phe | | Cys | Glu | | | Ser
380 | | Arg | Gly | Pro |
| Arg
385 | Met | ੋਪ੍ਰਾਤ | Arg | Cys | Leu
390 | Arg | Glu | Glu | Asp | Arg
395 | Ser | Leu | Asn | Gln | Tyr
400 |
| Pro | Ala | Leu | Tyr | Tyr
405 | Pro | Glu | Leu | Tyr | Ile
410 | Leu | Lys | Gly | Gly | Tyr
415 | Arg |
| Asp | Phe | Phe | Pro
420 | Glu | Tyr | Met | Glu | Leu
425 | Cys | Glu | Pro | Gln | Ser
430 | Tyr | Cys |
| | | | | | | | | | . 7 | | | | | | |

Leu Leu Val Lys Asp Met Ser Pro 465 470

<210> 1902

<211> 476

<212> PRT

<213> Homo sapiens

<400> 1902

Met Ala Val Pro Phe Val Glu Asp Trp Asp Leu Val Gln Thr Leu Gly
1 10 15

Glu Gly Ala Tyr Gly Glu Val Gln Leu Ala Val Asn Arg Val Thr Glu 20 25 30

Glu Ala Val Ala Val Lys Ile Val Asp Met Lys Arg Ala Val Asp Cys 35 40 45

Pro Glu Asn Ile Lys Lys Glu Ile Cys Ile Asn Lys Met Leu Asn His 50 55 60

Glu Asn Val Val Lys Phe Tyr Gly His Arg Arg Glu Gly Asn Ile Gln 65 70 75 80

Tyr Leu Phe Leu Glu Tyr Cys Ser Gly Gly Glu Leu Phe Asp Arg Ile 85 90 95

Glu Pro Asp Ile Gly Met Pro Glu Pro Asp Ala Gln Arg Phe Phe His 100 105 110

Gln Leu Met Ala Gly Val Val Tyr Leu His Gly Ile Gly Ile Thr His 115 120 125

Arg Asp Ile Lys Pro Glu Asn Leu Leu Asp Glu Arg Asp Asn Leu 130 135 140

Lys Ile Ser Asp Phe Gly Leu Ala Thr Val Phe Arg Tyr Asn Asn Arg 145 150 155 160

Glu Arg Leu Leu Asn Lys Met Cys Gly Thr Leu Pro Tyr Val Ala Pro 165 170 175

Glu Leu Leu Lys Arg Arg Glu Phe His Ala Glu Pro Val Asp Val Trp
180 185 190

Ser Cys Gly Ile Val Leu Thr Ala Met Leu Ala Gly Glu Leu Pro Trp 195 200 205

Asp Gln Pro Ser Asp Ser Cys Gln Glu Tyr Ser Asp Trp Lys Glu Lys 210 215 220

Pro Asp Ile Lys Lys Asp Arg Trp Tyr Asn Lys Pro Leu Lys Lys Gly 260 265 270

Ala Lys Arg Pro Arg Val Thr Ser Gly Gly Val Ser Glu Ser Pro Ser 275 280 285

Gly Phe Ser Lys His Ile Gln Ser Asn Leu Asp Phe Ser Pro Val Asn 290 295 300

Ser Ala Ser Ser Glu Glu Asn Val Lys Tyr Ser Ser Ser Gln Pro Glu 305 310 315 320

Pro Arg Thr Gly Leu Ser Leu Trp Asp Thr Ser Pro Ser Tyr Ile Asp 325 330 335

Lys Leu Val Gln Gly Ile Ser Phe Ser Gln Pro Thr Cys Pro Asp His $340 \ \ 345 \ \ 350$

Met Leu Leu Asn Ser Gln Leu Leu Gly Thr Pro Gly Ser Ser Gln Asn 355 360 365

Pro Trp Gln Arg Leu Val Lys Arg Met Thr Arg Phe Phe Thr Lys Leu 370 375 380

Asp Ala Asp Lys Ser Tyr Gln Cys Leu Lys Glu Thr Cys Glu Lys Leu 385 390 395 400

Gly Tyr Gln Trp Lys Lys Ser Cys Met Asn Gln Val Thr Ile Ser Thr 405 410 415

Thr Asp Arg Arg Asn Asn Lys Leu Ile Phe Lys Val Asn Leu Leu Glu 420 425 430

Met Asp Asp Lys Ile Leu Val Asp Phe Arg Leu Ser Lys Gly Asp Gly 435 440 445

Leu Glu Phe Lys Arg His Phe Leu Lys Ile Lys Gly Lys Leu Ile Asp
450 460

Ile Val Ser Ser Gln Lys Val Trp Leu Pro Ala Thr 465 470 475

<210> 1903

<211> 542

<212> PRT

<213> Homo sapiens

<400> 1903

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and the second of the second o

- Ser Gln Ser Ser His Ser Ser Ser Gly Thr Leu Ser Ser Leu Glu Thr 50 55 60
- Val Ser Thr Gln Glu Leu Tyr Ser Ile Pro Glu Asp Gln Glu Pro Glu 65 70 75 80
- Asp Gln Glu Pro Glu Glu Pro Thr Pro Ala Pro Trp Ala Arg Leu Trp 85 90 95
- Ala Leu Gl
n Asp Gly Phe Ala Asn Leu Glu Cys Val Asn Asp Asn Tyr
 100 \$100\$
- Trp Phe Gly Arg Asp Lys Ser Cys Glu Tyr Cys Phe Asp Glu Pro Leu 115 120 125
- Leu Lys Arg Thr Asp Lys Tyr Arg Thr Tyr Ser Lys His Phe Arg 130 135 140
- Asp His Ser Gly Asn Gly Thr Phe Val Asn Thr Glu Leu Val Gly Lys 165 170 175
- Gly Lys Arg Arg Pro Leu Asn Asn Ser Glu Ile Ala Leu Ser Leu 180 185 190
- Ser Arg Asn Lys Val Phe Val Phe Phe Asp Leu Thr Val Asp Asp Gln 195 200 205
- Ser Val Tyr Pro Lys Ala Leu Arg Asp Glu Tyr Ile Met Ser Lys Thr 210 215 220
- Leu Gly Ser Gly Ala Cys Gly Glu Val Lys Leu Ala Phe Glu Arg Lys 225 230 235 240
- Thr Cys Lys Val Ala Ile Lys Ile Ile Ser Lys Arg Lys Phe Ala 245 250 255
- Ile Gly Ser Ala Arg Glu Ala Asp Pro Ala Leu Asn Val Glu Thr Glu 260 265 270
- Ile Glu Ile Leu Lys Lys Leu Asn His Pro Cys I.e Ile Lys 1le Lys 275 280 285
- Asn Phe Phe Asp Ala Glu Asp Tyr Tyr Ile Val Leu Glu Leu Met Glu 290 295 300
- Gly Glu Leu Phe Asp Lys Val Val Gly Asn Lys Arg Leu Lys Glu 305 310 315 320
- Ala Thr Cys Lys Leu Tyr Phe Tyr Gln Met Leu Leu Ala Val Gln Tyr

Leu Leu Ser Ser Gln Glu Glu Asp Cys Leu Ile Lys Ile Thr Asp Phe 355 360 365

Gly His Ser Lys Ile Leu Gly Glu Thr Ser Leu Met Arg Thr Leu Cys 370 375 380

Gly Thr Pro Thr Tyr Leu Ala Pro Glu Val Leu Val Ser Val Gly Thr 385 390 395 400

Ala Gly Tyr Asn Arg Ala Val Asp Cys Trp Ser Leu Gly Val Ile Leu 405 410 415

Phe Ile Cys Leu Ser Gly Tyr Pro Pro Phe Ser Glu His Arg Thr Gln 420 425 430

Val Ser Leu Lys Asp Gln Ile Thr Ser Gly Lys Tyr Asn Phe Ile Pro 435 440 445

Glu Val Trp Ala Glu Val Ser Glu Lys Ala Leu Asp Leu Val Lys Lys 450 455 460

Leu Leu Val Val Asp Pro Lys Ala Arg Phe Thr Thr Glu Glu Ala Leu 465 470 475 480

Arg His Pro Trp Leu Gln Asp Glu Asp Met Lys Arg Lys Phe Gln Asp 485 490 495

Leu Leu Ser Glu Glu Asn Glu Ser Thr Ala Leu Pro Gln Val Leu Ala 500 505 510

Gln Pro Ser Thr Ser Arg Lys Arg Pro Arg Glu Gly Glu Ala Glu Gly
515 520 525

Ala Glu Thr Thr Lys Arg Pro Ala Val Cys Ala Ala Val Leu 530 535 540

<210> 1904

<211> 12

<212> PRT

<213> Artificial Sequence

2020s

<400> 1904

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<210> 1905

<211> 9

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peptide
<400> 1905
Gly Gly Arg Ser Pro Ala Met Pro Glu
<210> 1906
<211> 12
<212> PRT
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<400> 1906
Leu Tyr Arg Ser Pro Ser Ala Met Pro Glu Asn Leu
<210> 1907
<211> 11
<212 > PRT
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<222> (8)
<223> Pro, Phe, Tyr or Trp
<400> 1907
Leu Tyr Arg Ser Pro Ser His Xaa Glu Asn Leu
<210> 1908
<211> 11
<2112> PRT
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<2230 Description of Artificial Sequence: Synthetic
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<220>
<221> MOD RES
<222: (8)
<223> Pro, Phe, Tyr or Trp
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<210> 1909
<211> 11
<212> PRT
<213> Artificial Sequence
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<220>
<221> MOD RES
<222> (8)
<223> Pro, Phe, Tyr or Trp
<400> 1909
Leu Tyr Thr Ser Pro Ser Tyr Xaa Glu Asn Leu
<210> 1910
<211> 11
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      peptide
<220>
<221> MOD RES
<222> (8)
<223> Pro, Phe, Tyr or Trp
<400> 1910
Leu Tyr Thr Ser Pro Ser His Xaa Glu Asn Leu
<210> 1911
<211> 11
<212> PRT
<113> Artificial Sequence
<120>
<223> Description of Artificial Sequence: Synthetic
      peptide
<220>
<221> MOD RES
<222> (8)
<223> Pro, Phe, Tyr or Trp
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<210> 1912
<211> 11
<212> PRT
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<220>
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<223> Pro, Phe, Tyr or Trp
<400> 1912
Trp Tyr Arg Ser Pro Ser Phe Xaa Glu Asn Leu
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<220>
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<223> Pro, Phe, Tyr or Trp
<400> 1913
Trp Tyr Thr Ser Pro Ser His Xaa Glu Asn Leu
<210> 1914
<211> 11
<212> PRT
<213> Artificial Sequence
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<220>
<221> MOD RES
<222> (8)
<223> Pro, Phe, Tyr or Trp
<4(C> 1914
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<211> 11
<212> PRT
<213> Artificial Sequence
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<220>
<221> MOD RES
<222> (8)
<223> Pro, Phe, Tyr or Trp
<400> 1915
Phe Tyr Ser Ser Pro Ser His Xaa Glu Asn Leu
<210> 1916
<211> 11
<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic
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<220>
<221> MOD RES
<222> (8)
<223> Pro, Phe, Tyr or Trp
<400> 1916
Phe Tyr Thr Ser Pro Ser Met Xaa Glu Asn Leu
<210> 1917
<211> 11
<212> PRT
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<11.05
<221> MOD RES
<222> (8)
<223> Pro, Phe, Tyr or Trp
<400> 1917
Phe Tyr Thr Ser Pro Ser Phe Xaa Glu Asn Leu
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<223> Description of Artificial Sequence: Synthetic
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<222> (8)
<223> Pro, Phe, Tyr or Trp
<400> 1918
Phe Tyr Thr Ser Pro Ser Tyr Xaa Glu Asn Leu
<210> 1919
<211> 11
<212> PRT
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<220>
<221> MOD RES
<222> (8)
<223> Pro, Phe, Tyr or Trp
<400> 1919
Trp Tyr Thr Ser Pro Ser Met Xaa Glu Asn Leu
<210> 1920
<211> 11
<212> PRT
<213> Artificial Sequence
<113> Description of Artificial Sequence: Synthetic
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<1110>
<111> MOD RES
<222> (8)
<223> Pro, Phe, Tyr or Trp
Trp Tyr Thr Ser Pro Ser Phe Xaa Glu Asn Leu
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<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic
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<221> MOD RES
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<223> Pro, Phe, Tyr or Trp
<400> 1921
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<210> 1922
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<212> PRT
<213> Artificial Sequence
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<400> 1922
Leu Ala Arg Ser Ala Ser Met Pro Glu Ala Leu
1
           5
<210> 1923
<211> 11
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
     peptide
<400> 1923
Arg Tyr Ser Leu Pro Pro Glu Leu Ser Asn Met
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<2:10> 1924
<211> 11
<112> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
     pertide
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<210> 1925
<211> 11
<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic
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<400> 1925
Leu Tyr Arg Ser Pro Ser Met Pro Glu Asn Leu
            5
<210> 1926
<211> 11
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<400> 1930
Tyr Xaa Xaa Pro Ser Xaa Xaa Xaa Asn
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<4) +> 1931
Gly Gly Ser Pro Ser Met
<210> 1932
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Ser Met Pro Glu Asn Leu
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<210> 1933
<211> 22
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Tyr Gly Arg Lys Lys Arg Arg Gln Arg Arg Arg Leu Tyr Arg Ser Pro
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Ala Met Pro Glu
<210> 1935
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<210> 1937
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Pro Glu Leu Ser Asn Met
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<210> 1939
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Tyr Gly Arg Lys Lys Arg Arg Gln Arg Arg Gly Gly Ser Pro Ala
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Met Pro
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<210> 1941
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<400> 1942
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<210> 1943
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<400> 1943
Tyr Gly Arg Lys Lys Arg Arg Gln Arg Arg Arg Gly Gly Ser Pro Ser
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Met
<210> 1944
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<400> 1944
Tyr Gly Arg Lys Lys Arg Arg Gln Arg Arg Arg Tyr Gly Gly Gly Gly
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<210> 1945
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Tyr Gly Arg Lys Lys Arg Arg Gln Arg Arg Arg Tyr Gly Gly Gly Gly
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Gly Gly Asn
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 Pro Asp Tyr Ala
              20
<210> 1948
 <211> 22
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 Asp Asp Asp Lys Gly
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